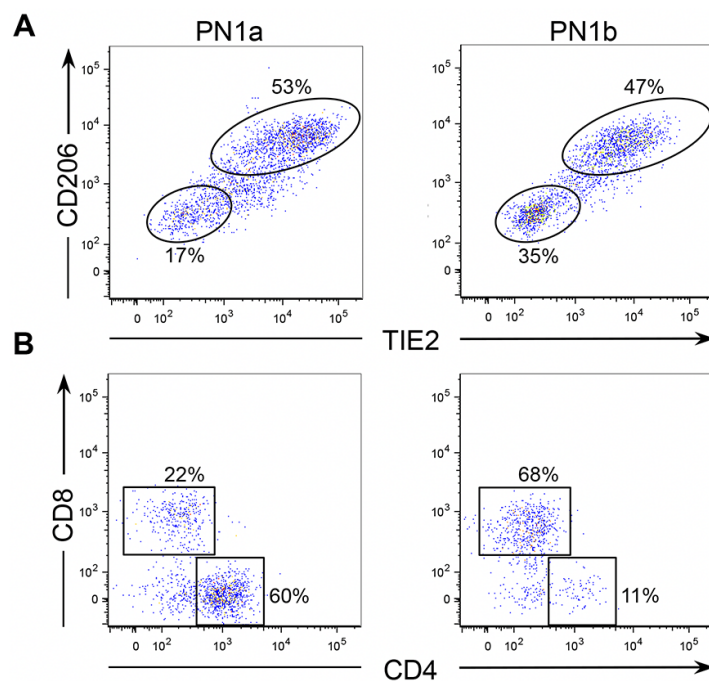
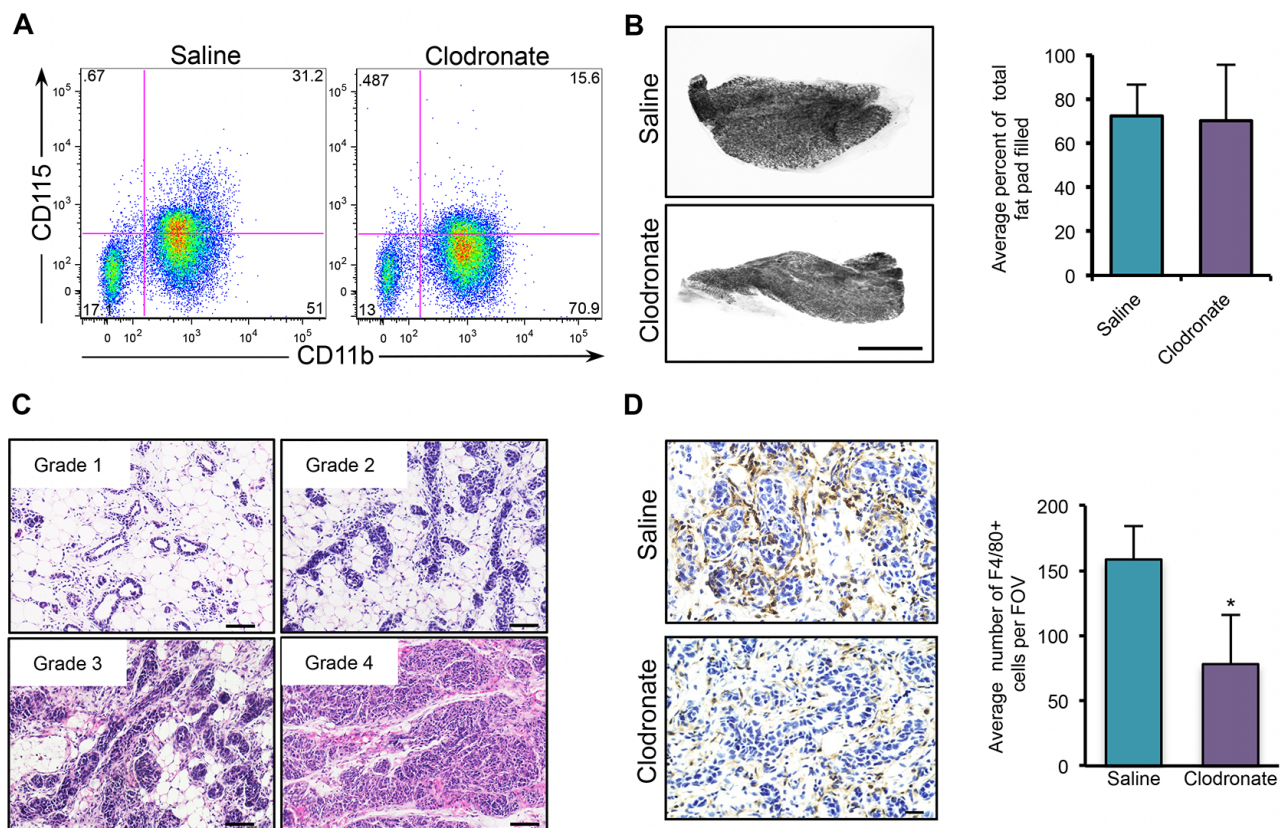


Macrophages promote the progression of premalignant mammary lesions to invasive cancer

SUPPLEMENTARY MATERIALS



Supplementary Figure 1: Myeloid and lymphoid populations in PN1a and PN1b lesions. **A.** Dot plots show CD206 and TIE2-expressing macrophages after gating on CD45⁺CD11b⁺F4/80⁺ cells in PN1a and PN1b lesions at 16 weeks post-transplantation. **B.** Dot plots depict the proportion of CD8⁺ (cytotoxic T cells) and CD4⁺ (T helper cells) lymphocytes after gating on CD45⁺CD11b⁺CD3⁺ cells in PN1a and PN1b lesions at 16 weeks post-transplantation. Data shown represent 1 experiment using 2 pooled lesions (2 mice) in each group, and the experiments were repeated 2-4 times.



Supplementary Figure 2: Macrophage depletion in PN1a-bearing mice. **A.** Dot blot depicts the number of CD11b⁺CD115⁺ cells that were isolated from the bone marrow of saline or clodronate liposome-treated mice bearing PN1a lesions. Cells were gated on SYTOX red⁻ and CD45⁺ subpopulations, and a minimum of 3 mice (3 lesions) were analyzed per a group. **B.** Representative images of carmine-stained saline- and clodronate liposome-treated PN1a lesions. scale bar = 2.5 mm (left). Graph depicts the percent of total fat pad filled where a minimum of 10 lesions (6 mice) were analyzed per group. Values are mean + SD, $p=0.79$. **C.** H&E staining demonstrating histological grades of PN1a lesions. Grade 1 lesions are characterized by diffuse, well-organized glandular patterns with a single layer of luminal epithelial cells surrounding a central lumen with focal or multifocal hyperplastic regions. Grade 2 lesions have diffuse, well-differentiated hyperplastic ductal and lobuloalveolar patterns, while grade 3 lesions show multifocal regions consisting of solid nests and hyperplastic ductal structures with cytologic atypia. Grade 4 lesions are characterized by solid nests of epithelial cells with little or no glandular differentiation and cytologic atypia. Scale bars = 10 μ m. **D.** Representative images of saline- and clodronate liposome-treated lesions stained with an antibody to F4/80 to detect macrophages (left). Graph represents the average number of F4/80⁺ cells per field of view (FOV). Ten FOV were counted for each lesion at 20X magnification, and a minimum of a 10 lesions (6 mice) were analyzed for each group. Values are mean + SD. * $p<0.001$, Scale bar = 10 μ m.

Supplementary Table 1: List of antibodies for immunostaining and flow cytometry

Antibody	Application	Manufacturer	Clone	Dilution
CK8	IF	DSHB	TROMA-1	1:250
CK8	IF	Biologend	Poly19053	1:200
CK14	IF	Covance	PRB-155P	1:400
CK8	IF	Progen Biotechnik	18.04	1:200
pan-CK	IF	Abcam	AE1/AE3+5D3	1:50
integrin $\alpha 6$	IF	BD Biosciences	GoH3	1:200
Ki67	IF	Abcam	SP6	1:100
Laminin	IF	Sigma	L9393	1:100
F4/80	IHC	AbD Serotec	A3-1	1:100
F4/80	FC	Biologend	BM8	1:100
MHCII	FC	Biologend	I-A/I-E	1:100
TIE2	FC	Biologend	TEK4	1:200
CD3	FC	Biologend	145-2C11	1:200
CD4	FC	Biologend	GK1.5	1:200
CD8	FC	Biologend	53-6.7	1:200
CD11b	FC	eBioscience	M1/70	1:100
CD45	FC	BD Biosciences	30-F11	1:100
CD204	FC	BD Biosciences	2F8	1:100
CD206	FC	Biologend	C068C2	1:100
CD206	FC	Abcam	EPR6828(B)	1:200

IF: immunofluorescence; IHC: immunohistochemistry; FC: flow cytometry.

Supplementary Table 2: Complete list of differential gene expression in PN1a, PN1b and p53-null mammary glands.

See Supplementary File 1

Supplementary Table 3: Primer sequences for qPCR

Gene symbol	Sense primer (5'-3')	Antisense primer (5'-3')
<i>Il6</i>	AGTCAATTCCAGAAACCGCTATGA	TAGGGAAGGCCGTGGTTGT
<i>Il10</i>	CAGAGCCACATGCTCCTAGA	TGTCCAGCTGGTCCTTTGTT
<i>Il12p40</i>	CAGCCGAGTGATGTACAAGG	TAAACGGGAAATCTGCACCT
<i>Arg1</i>	TTCTCAAAGGACAGCCTCG	CAGACCGTGGGTTCCTCACA
<i>Nos2</i>	GTCAACTGCAAGAGAACGGAGA	CTGAGAACAGCACAAGGGGTT
<i>Vegfa</i>	AGGCTGCTGTAACGATGAAG	TCTCCTATGTGCTGGCTTTG
<i>Tgfb</i>	TGGAGCAACATGTGGAATC	GTCAGCAGCCGGTTACCA
<i>Tnfa</i>	CTGTAGCCACGTCGTAGC	TTGAGATCCATGCCGTTG
<i>18s</i>	GTAACCCGTTGAACCCCAT	CCATCCAATCGGTAGTAGCG
<i>Gas6</i>	GGATTTGCTACCTACAGGCTCA	TAACTTCCCAGGTGGTTTCC
<i>Gapdh</i>	GCTACACTGAGGACCAGTTGT	CTCCTGTTATTATGGGGGTCTG

^a Sequences were designed using the Universal Probe Library Assay Design Center, Roche Applied Biosciences (<http://qpcr.probefinder.com/roche3.html>)